

CONFIDENTIAL

EXHIBIT 1

EXHIBIT 1

PSEUDO CODE LISTING OF FOLLOW UP FOR ASTHMA OFFICE VISIT DIALOG

```

1. package com.solvech.st_notify;
2. import java.util.*;
3. import com.solvech.st_lang.*;
4. public class FU_Asthma_Dialog extends DialogBase
5. {
6.     private static final String ATTR_DOCTOR_USES_PFR1      = "person(doctor).uses_pfr1";
7.     private static final String ATTR_DOCTOR_USES_PFR2      = "person(doctor).uses_pfr2";
8.
9.     // is the inOfficePF an attribute of the patient or of the dialog??
10.    private static final String ATTR_PATIENT_INOFFICEPF      = "person(patient).inOfficePF";
11.
12.    private static final String ATTR_PATIENT_AGE = "person(patient).age";
13.    private static final String ATTR_PATIENT_HEIGHT = "person(patient).height";
14.    private static final String ATTR_PATIENT_SEX = "person(patient).sex";
15.
16.    private final static String ATTR_DIALOG_WORSE           = "dialog.worse";
17.    private final static String ATTR_DIALOG_READY           = "dialog.ready";
18.    private final static String ATTR_DIALOG_RETRIG          = "dialog.retrig";
19.    private final static String ATTR_DIALOG_RECALL_1        = "dialog.recall_1";
20.    private final static String ATTR_DIALOG_RECALL_2        = "dialog.recall_2";
21.    private final static String ATTR_DIALOG_OK_END          = "dialog.okend";

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19. private final static String ATTR_DIALOG_ACK_ER      = "dialog.ackER";
20. private final static String ATTR_DIALOG_PF_1      = "dialog.pf_1";

21. //=====
22. private boolean doctor_uses_pfr1;
23. private boolean doctor_uses_pfr2;
24. private int inOfficePF;
25. private int retrig;
26. private int age = 0;
27. private int height = 0;
28. private String sex = "";
29.
30. public FU_Asthma_Dialog()
31. {
32.     super();
33. }

34. final protected void init()
35.     throws Exception
36. {
37.     // load up all the variables passed in from the message.
38.     doctor_uses_pfr1 = getAttrBoolean(ATTR_DOCTOR_USES_PFR1);
39.     doctor_uses_pfr2 = getAttrBoolean(ATTR_DOCTOR_USES_PFR2);
40.     inOfficePF      = getAttrInt(ATTR_PATIENT_INOFFICEPF);
41.     retrig          = getAttrInt(ATTR_DIALOG_RETRIG);
42.     age             = getAttrInt(ATTR_PATIENT_AGE);

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= getAttrInt(ATTR_PATIENT_HEIGHT);
= getAttrString(ATTR_PATIENT_SEX);

```

```

height
sex

```

```

}

```

```

final protected void register()
{

```

```

// registration is a way of telling scheduler what attributes are

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// needed for a message.

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```

registerAttribute(ATTR_DOCTOR_USES_PFR1);
registerAttribute(ATTR_DOCTOR_USES_PFR2);
registerAttribute(ATTR_PATIENT_INOFFICEPF);
registerAttribute(ATTR_PATIENT_AGE);
registerAttribute(ATTR_PATIENT_HEIGHT);
registerAttribute(ATTR_PATIENT_SEX);
registerAttribute(ATTR_DIALOG_WORSE);
registerAttribute(ATTR_DIALOG_READY);
registerAttribute(ATTR_DIALOG_RETRIG);
registerAttribute(ATTR_DIALOG_RECALL_1);
registerAttribute(ATTR_DIALOG_RECALL_2);
registerAttribute(ATTR_DIALOG_OK_END);
registerAttribute(ATTR_DIALOG_ACK_ER);
registerAttribute(ATTR_DIALOG_PF_1);

```

```

// register the Prompts that we are using. So that
// they can be verified before the conversation begins.
registerPrompt(PROMPT_ID_GOTO_ER);

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67. registerPrompt(PROMPT_ID_OK_END);
68. registerPrompt(PROMPT_ID_PEAK_FLOW);
69. registerPrompt(PROMPT_ID_READY_PF1);
70. registerPrompt(PROMPT_ID_RECALL_30);
71. registerPrompt(PROMPT_ID_WORSE);

72. registerPrompt(PROMPT_ID_ALERT_WORSE);
73. registerPrompt(PROMPT_ID_ALERT_NO_RESPONSE);
74. registerPrompt(PROMPT_ID_ALERT_PF1_BELOW_OFFICE_PF);
75. registerPrompt(PROMPT_ID_ALERT_PF1_BELOW_50_PERCENT_PREDICTED);
76. registerPrompt(PROMPT_ID_ALERT_STAT_CHECK);

77.
78. // register the roles that we require
79. registerRole(ROLE_PATIENT);
80. registerRole(ROLE_DOCTOR);
81. }

82. final protected void term()
83.     throws Exception
84.     {
85.     }

86. final protected void execute()
87.     {
88.         Phrase hello = new Phrase();
89.         hello.addMessage("Hello! Are you, ");

```

```

90. String patientName = getPatientName();
91. hello.addName(patientName);
92. hello.addMessage("? ");
93. int areYouTheOne = askYesNo(hello);
94. if(areYouTheOne != ASK_RESPONSE_YES)
95. {
96.     setStatus(DIALOG_STATUS_NORESPONSE);
97.     return;
98. }

99. Phrase messageFrom = new Phrase();
100. messageFrom.addMessage("There is a message from your doctor, ");
101. String doctorName = getDoctorName();
102. messageFrom.addName(doctorName);
103. messageFrom.addMessage(", recorded at");
104. messageFrom.addTime();
105. if(!getAcknowledgement(messageFrom))
106. {
107.     setStatus(DIALOG_STATUS_FAIL);
108.     return;
109. }

110. if(retrig <= 0)
111. {
112.     // [D1Q1] first question
113.     int worse = askYesNo(PROMPT_ID_WORSE);
114.

```

```

115. saveAnswer(ATTR_DIALOG_WORSE,worse);

116. if(worse == ASK_RESPONSE_YES)
117.     alertDoctor(PROMPT_ID_ALERT_WORSE);
118. else if(worse == ASK_RESPONSE_NONE)
119.     {
120.         alertDoctor(PROMPT_ID_ALERT_NO_RESPONSE);
121.         return;
122.     }
123. }

124. // [D1Q2] second question
125. int ready = askYesNo(PROMPT_ID_READY_PF1);
126. saveAnswer(ATTR_DIALOG_READY,ready);
127. if(ready != ASK_RESPONSE_YES)
128.     {
129.         retrig++;
130.         switch(retrig)
131.         {
132.             case 1:
133.                 {
134.                     saveAnswer(ATTR_DIALOG_RETRIG,retrig);
135.                     saveAnswer(ATTR_DIALOG_RECALL_1,getAcknowledgement(PROMPT_ID_RECALL_30));
136.                     callAgainMinutes(30);
137.                     break;

```

```

138.     }
139.     case 2:
140.     {
141.         saveAnswer(ATTR_DIALOG_RETRIG,retrig);
142.
143.         saveAnswer(ATTR_DIALOG_RECALL_2,getAcknowledgement(PROMPT_ID_RECALL_30));
144.         callAgain(30,TIME_UNITS_MINUTES);
145.         break;
146.     }
147.     case 3:
148.     {
149.         alertDoctor(PROMPT_ID_ALERT_NO_PF1);
150.         setStatus(DIALOG_STATUS_UNKNOWN);
151.         break;
152.     }
153.     return;
154. }
155. // [D1Q5] third question
156. int pf = askInt(PROMPT_ID_PEAK_FLOW);
157. saveAnswer(ATTR_DIALOG_PF_1,pf);
158. if(peakflow_rule1(pf) && doctor_uses_pfr1)
159. {
160.     alertDoctor(PROMPT_ID_ALERT_PF1_BELOW_OFFICE_PF);
161. }

```



```

162. else if(peakflow_rule2(pf) && doctor_uses_pfr2)
163. {
164.     alertDoctor(PROMPT_ID_ALERT_PF1_BELOW_50_PERCENT_PREDICTED);
165. }
166. else
167. {
168.     // [D1Q7] - ready to do dialog 2 on next scheduling time.
169.     saveAnswer(ATTR_DIALOG_OK_END,
170.         getAcknowledgement(PROMPT_ID_OK_END));
171.     return;
172. }
173. // [D1Q6]
174. saveAnswer(ATTR_DIALOG_ACK_ER, getAcknowledgement(PROMPT_ID_GOTO_ER));
175. alertDoctor(PROMPT_ID_ALERT_STAT_CHECK);
176. }
177. private boolean peakflow_rule1(int pf)
178. {
179.     if(pf < inOfficePF)
180.         return true;
181.     return false;
182. }
183. // compare peak flow to a formula to compute pf based on height, age, sex

```

```

184. private boolean peakflow_rule2(int pf)
185. {
186.     double predicted_pf = 0;

187.     if(sex.equalsIgnoreCase("male"))
188.     {
189.         predicted_pf = (((double)height * 0.0254 * 5.48) + 1.58) - ((double)age * 0.041)) *
60.0;
190.     }
191.     else
192.     {
193.         predicted_pf = (((double)height * 0.0254 * 3.72) + 2.24) - ((double)age * 0.03)) * 60.0;
194.     }
195.

196.     double percent = ((double)pf * 100.0) / predicted_pf;
197.     if(percent < 50.0)
198.         return true;

199.     return false;
200. }
201. }

```

EXHIBIT 2

EXHIBIT 2: PSEUDO CODE LISTING FOR CHEMISTRY BATTERY TEST RESULT DIALOG

```

1 package com.solveotech.st_notify;
2 import java.util.*; com.solveotech.st_lang.*;
3
4 public class ChemicalBatteryTest_Dialog extends DialogBase
5 {
6     private static final int COMPARE_TEST_NORMAL = 0;
7     private static final int COMPARE_TEST_LOW = -1;
8     private static final int COMPARE_TEST_HIGH = 1;
9     private static final int COMPARE_TEST_INDETERMINATE = -2;
10
11     // attributes
12     private static final String ATTR_PATIENT_AGE = "person(patient).age";
13     private static final String ATTR_PATIENT_SEX = "person(patient).sex";
14
15     // constants for test results
16     private static final int TEST_FIELD_NAME = 0;
17     private static final int TEST_FIELD_RESULT = 1;
18     private static final int TEST_FIELD_NORMAL_LOW = 2;
19     private static final int TEST_FIELD_NORMAL_HIGH = 3;
20     private static final int TEST_FIELD_UNITS = 4;
21     private static final int TEST_FIELD_REFERENCE_NORMAL = 5;
22     private static final int TEST_FIELD_REFERENCE_LOW = 6;
23     private static final int TEST_FIELD_REFERENCE_HIGH = 7;
24     private static final int TEST_FIELD_REFERENCE_INDETERMINATE = 8;
25
26     private static final int TEST_OUT_NAME = 0;
27     private static final int TEST_OUT_RESULT = 1;
28     private static final int TEST_OUT_NORMAL_LOW = 2;
29     private static final int TEST_OUT_NORMAL_HIGH = 3;
30

```

```

31 private static final int TEST_OUT_YOUR_RESULT = 4;
32 // computed - not sent
33 private static final int TEST_OUT_UNITS = 5;
34 private static final int TEST_OUT_REFERENCE = 6;
35 private static final int TEST_OUT_MAX = 7;
36
37 private static final int TEST_TERSE_OUT_NAME = 0;
38 private static final int TEST_TERSE_OUT_RESULT = 1;
39 private static final int TEST_TERSE_OUT_UNITS = 2;
40 private static final int TEST_TERSE_OUT_MAX = 3;
41
42 private static final String TEST_COLUMN_YOUR_RESULT = "yourResult";
43 private static final String TEST_COLUMN_REFERENCE = "reference";
44
45
46 // headers
47 // these must stay in sync with the TEST_FIELD_... constants defined above
48 private static String[] TEST_HEADER_NAMES =
49 {
50     "dialog.test.name",
51     "dialog.test.result",
52     "dialog.test.normal_low",
53     "dialog.test.normal_high",
54     "dialog.test.units",
55     "dialog.test.reference_normal",
56     "dialog.test.reference_low",
57     "dialog.test.reference_high",
58     "dialog.test.reference_indeterminate"
59 };
60
61

```

```

62 // indexes into array TEST_ATTR_FIELD_NAMES
63
64 private static final int TEST_INDEX_GLUCOSE
65 private static final int TEST_INDEX_UREA_NITROGEN
66 private static final int TEST_INDEX_CREATININE
67 private static final int TEST_INDEX_CALCIIUM
68 private static final int TEST_INDEX_SODIUM
69 private static final int TEST_INDEX_POTASSIUM
70 private static final int TEST_INDEX_CO2
71 private static final int TEST_INDEX_CHLORIDE
72 private static final int TEST_INDEX_TOTAL_PROTEIN
73 private static final int TEST_INDEX_ALBUMIN
74 private static final int TEST_INDEX_GLOBULIN
75 private static final int TEST_INDEX_A_G_RATIO
76 private static final int TEST_INDEX_TOTAL_BILIRUBIN
77 private static final int TEST_INDEX_ALKALINE_PHOSPHATASE
78 private static final int TEST_INDEX_AST_SGOT
79 private static final int TEST_INDEX_AST_SGPT
80
81
82 // Table of attribute names - this will be used to fetch
83 // the values of the attributes so that the results can be reported.
84 private static String[][] TEST_ATTR_FIELD_NAMES =
85 {
86     {
87         "dialog.test.name.glucose",
88         "dialog.test.result.glucose",
89         "dialog.test.normal_low.glucose",
90         "dialog.test.normal_high.glucose",
91         "dialog.test.units.glucose",
92         "dialog.test.reference_normal.glucose",
93         "dialog.test.reference_low.glucose",
94     }
95 }

```

```

93 "dialog.test.reference_high.glucose",
94 "dialog.test.reference_indeterminate.glucose"
95 },
96
97 {
98     "dialog.test.name.urea_nitrogen",
99     "dialog.test.result.urea_nitrogen",
100     "dialog.test.normal_low.urea_nitrogen",
101     "dialog.test.normal_high.urea_nitrogen",
102     "dialog.test.units.urea_nitrogen",
103     "dialog.test.reference_normal.nitrogen",
104     "dialog.test.reference_low.nitrogen",
105     "dialog.test.reference_high.nitrogen",
106     "dialog.test.reference_indeterminate.nitrogen"
107 },
108
109 {
110     "dialog.test.name.creatinine",
111     "dialog.test.result.creatinine",
112     "dialog.test.normal_low.creatinine",
113     "dialog.test.normal_high.creatinine",
114     "dialog.test.units.creatinine",
115     "dialog.test.reference_normal.creatinine",
116     "dialog.test.reference_low.creatinine",
117     "dialog.test.reference_high.creatinine",
118     "dialog.test.reference_indeterminate.creatinine"
119 },
120
121 {
122     "dialog.test.name.calcium",
123     "dialog.test.result.calcium",
124     "dialog.test.normal_low.calcium",
125     "dialog.test.normal_high.calcium",
126     "dialog.test.units.calcium",

```

```

124 "dialog.test.reference_normal.calcium",
125 "dialog.test.reference_low.calcium",
126 "dialog.test.reference_high.calcium",
127 "dialog.test.reference_indeterminate.calcium"
128 },
129 {
130 "dialog.test.name.sodium",
131 "dialog.test.result.sodium",
132 "dialog.test.normal_low.sodium",
133 "dialog.test.normal_high.sodium",
134 "dialog.test.units.sodium",
135 "dialog.test.reference_normal.sodium",
136 "dialog.test.reference_low.sodium",
137 "dialog.test.reference_high.sodium",
138 "dialog.test.reference_indeterminate.sodium"
139 },
140
141 {
142 "dialog.test.name.potassium",
143 "dialog.test.result.potassium",
144 "dialog.test.normal_low.potassium",
145 "dialog.test.normal_high.potassium",
146 "dialog.test.units.potassium",
147 "dialog.test.reference_normal.potassium",
148 "dialog.test.reference_low.potassium",
149 "dialog.test.reference_high.potassium",
150 "dialog.test.reference_indeterminate.potassium"
151 },
152
153 {
154 "dialog.test.name.co2",
155 "dialog.test.result.co2",

```



```

155 "dialog.test.normal_low.co2",
156 "dialog.test.normal_high.co2",
157 "dialog.test.units.co2",
158 "dialog.test.reference_normal.co2",
159 "dialog.test.reference_low.co2",
160 "dialog.test.reference_high.co2",
161 "dialog.test.reference_indeterminate.co2"
162 },
163
164 {
165 "dialog.test.name.chloride",
166 "dialog.test.result.chloride",
167 "dialog.test.normal_low.chloride",
168 "dialog.test.normal_high.chloride",
169 "dialog.test.units.chloride",
170 "dialog.test.reference_normal.chloride",
171 "dialog.test.reference_low.chloride",
172 "dialog.test.reference_high.chloride",
173 "dialog.test.reference_indeterminate.chloride"
174 },
175
176 {
177 "dialog.test.name.total_protein",
178 "dialog.test.result.total_protein",
179 "dialog.test.normal_low.total_protein",
180 "dialog.test.normal_high.total_protein",
181 "dialog.test.units.total_protein",
182 "dialog.test.reference_normal.total_protein",
183 "dialog.test.reference_low.total_protein",
184 "dialog.test.reference_high.total_protein",
185 "dialog.test.reference_indeterminate.total_protein"

```

```

186 },
187
188 {
189     "dialog.test.name.albumin",
190     "dialog.test.result.albumin",
191     "dialog.test.normal_low.albumin",
192     "dialog.test.normal_high.albumin",
193     "dialog.test.units.albumin",
194     "dialog.test.reference_normal.albumin",
195     "dialog.test.reference_low.albumin",
196     "dialog.test.reference_high.albumin",
197     "dialog.test.reference_indeterminate.albumin"
198 },
199
200 {
201     "dialog.test.name.globulin",
202     "dialog.test.result.globulin",
203     "dialog.test.normal_low.globulin",
204     "dialog.test.normal_high.globulin",
205     "dialog.test.units.globulin",
206     "dialog.test.reference_normal.globulin",
207     "dialog.test.reference_low.globulin",
208     "dialog.test.reference_high.globulin",
209     "dialog.test.reference_indeterminate.globulin"
210 },
211
212 {
213     "dialog.test.name.a_g_ratio",
214     "dialog.test.result.a_g_ratio",
215     "dialog.test.normal_low.a_g_ratio",
216     "dialog.test.normal_high.a_g_ratio",
217     "dialog.test.units.a_g_ratio",

```

```

217 "dialog.test.reference_normal.a_g_ratio",
218 "dialog.test.reference_low.a_g_ratio",
219 "dialog.test.reference_high.a_g_ratio",
220 "dialog.test.reference_indeterminate.a_g_ratio"
221 },
222
223 {
224 "dialog.test.name.total_bilirubin",
225 "dialog.test.result.total_bilirubin",
226 "dialog.test.normal_low.total_bilirubin",
227 "dialog.test.normal_high.total_bilirubin",
228 "dialog.test.units.total_bilirubin",
229 "dialog.test.reference_normal.total_bilirubin",
230 "dialog.test.reference_low.total_bilirubin",
231 "dialog.test.reference_high.total_bilirubin",
232 "dialog.test.reference_indeterminate.total_bilirubin"
233 },
234
235 {
236 "dialog.test.name.alkaline_phosphatase",
237 "dialog.test.result.alkaline_phosphatase",
238 "dialog.test.normal_low.alkaline_phosphatase",
239 "dialog.test.normal_high.alkaline_phosphatase",
240 "dialog.test.units.alkaline_phosphatase",
241 "dialog.test.reference_normal.alkaline_phosphatase",
242 "dialog.test.reference_low.alkaline_phosphatase",
243 "dialog.test.reference_high.alkaline_phosphatase",
244 "dialog.test.reference_indeterminate.alkaline_phosphatase"
245 },
246
247 {
248 "dialog.test.name.ast_sgot",

```

```

248 "dialog.test.result.ast_sgot",
249 "dialog.test.normal_low.ast_sgot",
250 "dialog.test.normal_high.ast_sgot",
251 "dialog.test.units.ast_sgot",
252 "dialog.test.reference_normal.ast_sgot",
253 "dialog.test.reference_low.ast_sgot",
254 "dialog.test.reference_high.ast_sgot",
255 "dialog.test.reference_indeterminate.ast_sgot"
256 },
257
258 {
259 "dialog.test.name.alt_sgpt",
260 "dialog.test.result.alt_sgpt",
261 "dialog.test.normal_low.alt_sgpt",
262 "dialog.test.normal_high.alt_sgpt",
263 "dialog.test.units.alt_sgpt",
264 "dialog.test.reference_normal.alt_sgpt",
265 "dialog.test.reference_low.alt_sgpt",
266 "dialog.test.reference_high.alt_sgpt",
267 "dialog.test.reference_indeterminate.alt_sgpt"
268 }
269 };
270
271
272

```

private static final String TEST_EXPLANATION =

```

"The Chemical Screening Battery consists of a group of tests of "+
"liver, kidney, and pancreas functioning as well as other blood tests" +
"that are important for health such as calcium, total protein, "+
"and albumin." + "\n" +
" The fasting blood glucose can indicate how well the pancreas is "+
"producing insulin; high levels are associated with diabetes... ";

```

```

279 //
280
281 // The following two tables contain explanations for abnormal results for
282 // each test. The first table applies to high test results; the second
283 // to lower than normal results.
284 // It must be kept in the same order as the TEST_INDEX constants
285 // defined above.
286
287 private static final String[] ABNORMAL_HIGH_TEST_EXPLANATION =
288 {
289     // TEST_INDEX_HIGH_GLUCOSE
290     "Elevated blood glucose can be the result of Diabetes Mellitus. If this has not been noted before, you should
291     discuss this result with your medical provider",
292
293     // TEST_INDEX_HIGH_UREA_NITROGEN
294     "High urea nitrogen can be associated with dehydration, kidney problems, or the use of certain medicines",
295
296
297     // TEST_INDEX_HIGH_CREATININE
298     "High creatinine can be associated with dehydration, kidney problems, or the use of certain medicines",
299
300     // TEST_INDEX_HIGH_CALCIIUM
301     "Elevated calcium can be associated with parathyroid gland disorders, certain medications and use of excess
302     vitamin D, kidney problems, and other disorders",
303     // TEST_INDEX_HIGH_SODIUM
304     "High sodium levels can result from dehydration, diarrhea, excess salt intake, and other disorders",
305
306     // TEST_INDEX_HIGH_POTASSIUM
307     "High potassium levels can come from excess potassium ingestion, renal problems, certain medications, and other
308     disorders",
309

```

310 // TEST_INDEX_HIGH_CO2
 311 "Elevated carbon dioxide levels can come from repeated vomiting, emphysema, and certain medications",
 312
 313
 314
 315 // TEST_INDEX_HIGH_CHLORIDE
 316 "Elevated chloride levels may occur with dehydration, parathyroid disease, adrenal disease, and certain metabolic
 317 disorders",
 318
 319 // TEST_INDEX_HIGH_TOTAL_PROTEIN
 320 "High protein levels can result from dehydration, liver disease, chronic infections, and chronic inflammatory
 321 conditions",
 322
 323 // TEST_INDEX_HIGH_ALBUMIN
 324 "Albumin values can be elevated in dehydration states",
 325
 326 // TEST_INDEX_HIGH_GLOBULIN
 327 "High globulin levels can occur in chronic infections, liver disease, or auto-immune diseases",
 328
 329 // TEST_INDEX_A_G_HIGH_RATIO
 330 "Please see comments regarding albumin and globulin",
 331
 332 // TEST_INDEX_HIGH_TOTAL_BILIRUBIN
 333 "Elevated bilirubin levels can result from liver disease or breakdown of red blood cells",
 334
 335
 336 // TEST_INDEX_HIGH_ALKALINE_PHOSPHATASE
 337 "Elevated levels of alkaline phosphatase occur with liver problems, gall bladder disease, and infectious
 338 mononucleosis",
 339
 340 // TEST_INDEX_HIGH_AST_SGOT

```

341 "Elevated AST levels occur in liver disease and infectious
342 mononucleosis",
343 // TEST_INDEX_HIGH_ALT_SGPT
344 "Increased ALT levels can occur in liver or pancreatic disease, heart or muscle disease, and infectious
345 mononucleosis "
346 };
347
348 // These entries must be kept in the same order as the
349 // TEST_INDEX constants defined above.
350
351 private static final String[] ABNORMAL_LOW_TEST_EXPLANATION =
352 {
353 // TEST_INDEX_LOW_GLUCOSE
354 "Low blood glucose can be caused by rare pancreatic or metabolic disorders or certain medications",
355
356 // TEST_INDEX_LOW_UREA_NITROGEN
357 "Low urea nitrogen can be associated with poor protein intake or excess fluid intake ",
358
359 // TEST_INDEX_LOW_CREATININE
360 "Low creatinine can be associated with decrease muscle mass",
361
362 // TEST_INDEX_LOW_CALCIIUM
363 "Low calcium levels can be associated with parathyroid gland disorders, vitamin D deficiency, kidney problems,
364 and other disorders",
365
366 // TEST_INDEX_LOW_SODIUM
367 "Low sodium levels can result from over hydration, diarrhea, diuretic and other medications, and other disorders",
368
369
370
371

```

372 // TEST_INDEX_LOW_POTASSIUM
 373 "Low potassium levels can result from adrenal disorders, certain medications such as diuretics, diabetes, and other
 374 disorders",
 375
 376 // TEST_INDEX_LOW_CO2
 377 "Low carbon dioxide levels can result from diarrhea, kidney problems, diabetes, and certain medications",
 378
 379 // TEST_INDEX_LOW_CHLORIDE
 380 "Low chloride levels can result from vomiting, diabetes, decreased adrenal function, and certain medications",
 381
 382 // TEST_INDEX_LOW_TOTAL_PROTEIN
 383 "Low protein levels can result from liver disease, malabsorption, and poor nutrition",
 384
 385 // TEST_INDEX_LOW_ALBUMIN
 386 "Low albumin levels can result from liver disease, malabsorption conditions, and poor nutrition",
 387
 388 // TEST_INDEX_LOW_GLOBULIN
 389 "Low globulin levels can occur with renal disease and certain congenital or acquired immune disorders",
 390
 391 // TEST_INDEX_A_G_LOW_RATIO
 392 "Low A/G ratios can result from liver disease, collagen disease, chronic infections, and poor nutrition",
 393
 394 // TEST_INDEX_LOW_TOTAL_BILIRUBIN
 395 "This is a liver test; low levels of bilirubin are not worrisome",
 396
 397
 398 // TEST_INDEX_LOW_ALKALINE_PHOSPHATASE
 399 "Low levels of alkaline phosphatase can occur with poor nutrition, hypothyroidism, severe anemia, and
 400 hypophosphatasia",
 401
 402


```

403 // TEST_INDEX_LOW_AST_SGOT
404 "This is a liver function test; low levels of AST are not worrisome",
405
406 // TEST_INDEX_LOW_ALT_SGPT
407 " This is a liver function test; low levels of ALT are not worrisome"
408 };
409
410 //=====
411 // the header names will contain the names of each column
412 // in the table of test results.
413 private String[] headerNames;
414
415 // the testRows contain the data values corresponding to the
416 // array of attribute constants, TEST_ATTR_FIELD_NAMES, defined above
417 // each row represents a different test.
418 // for instance testRows[3] would represent the values for the calcium
419 // test. The index, TEST_INDEX_CALCIIUM, would be used to address
420 // this row, ie. testRows[TEST_INDEX_CALCIIUM]. The values of a row
421 // would correspond (in order) to the test name, result, normal_low, normal_high,
422 // units and a reference. For instance testRows[TEST_INDEX_GLUCOSE][TEST_FIELD_RESULT]
423 // would contain the patients test result value for the glucose test.
424
425 private String[][] testRows;
426
427 // the results and normals are separated out so that they can be
428 // used in calculations. They are loaded up from the attribute information.
429 private double[] testResult;
430 private double[] testNormalLow;
431 private double[] testNormalHigh;
432
433 private String sex = "";

```

```

434 private int age = 0;
435
436 public ChemicalBatteryTest_Dialog()
437 {
438     super();
439 }
440
441 final protected void init()
442     throws Exception
443 {
444     headerNames = new String[TEST_HEADER_NAMES.length];
445     testRows = new String[TEST_ATTR_FIELD_NAMES.length][];
446
447     // load up all the variables passed in from the message.
448     // load the headers
449
450     for(int i=0;i<TEST_HEADER_NAMES.length;i++)
451     {
452         headerNames[i] = getAttrString(TEST_HEADER_NAMES[i]);
453     }
454
455     for(int i=0;i<TEST_ATTR_FIELD_NAMES.length;i++)
456     {
457         String[] row = TEST_ATTR_FIELD_NAMES[i];
458         testRows[i] = new String[row.length];
459
460         for(int j=0;j<testRows[i].length;j++)
461         {
462             testRows[i][j] = getAttrString(TEST_ATTR_FIELD_NAMES[i][j]);
463         }
464         testResult[i]= getAttrDouble(TEST_ATTR_FIELD_NAMES[i][TEST_FIELD_RESULT]);

```

```

465         testNormalLow[i] =
466         getAttrDouble(TEST_ATTR_FIELD_NAMES[i][TEST_FIELD_NORMAL_LOW]);
467         testNormalHigh[i] =
468         getAttrDouble(TEST_ATTR_FIELD_NAMES[i][TEST_FIELD_NORMAL_HIGH]);
469     }
470
471     age = getAttrInt(ATTR_PATIENT_AGE);
472     sex = getAttrString(ATTR_PATIENT_SEX);
473
474
475     final protected void register()
476     {
477         // registration is a way of telling scheduler what attributes are
478         // needed for a message.
479
480         // only the headers are registered
481         // It is quite possible that the patient only has
482         // had a subset of the tests. Registering
483         // all tests would result in the dialog aborting
484         // if any tests were missing.
485         for(int i=0;i<TEST_HEADER_NAMES.length;i++)
486         {
487             registerAttribute(TEST_HEADER_NAMES[i]);
488         }
489
490         // to compute the normal for the AKALINE_PHOSPHATASE test
491         // we need the age and sex of the patient.
492         registerAttribute(ATTR_PATIENT_AGE);
493         registerAttribute(ATTR_PATIENT_SEX);
494
495         // register the prompts that we are using. So that

```

```

496 // they can be verified before the conversation begins.
497 //
498 //
499 // register the roles that we require
500 registerRole(ROLE_PATIENT);
501 registerRole(ROLE_DOCTOR);
502 }
503
504 final protected void term()
505     throws Exception
506 {
507 }
508
509 final protected void execute()
510 {
511     // login has already been done.
512
513     Phrase messageFrom = new Phrase();
514     messageFrom.addMessage("Here are the results of your lab tests.");
515     if(!getAcknowledgement(messageFrom))
516     {
517         setStatus(DIALOG_STATUS_FAIL);
518         return;
519     }
520
521     Phrase abnormalHighTestExplanation = new Phrase();
522     Phrase abnormalLowTestExplanation = new Phrase();
523     boolean haveAbnormalTests = false;
524     boolean haveAbnormalLowTests = false;
525     boolean haveAbnormalHighTests = false;
526

```

```

527         for(int i=0;i<testRows.length;i++)
528         {
529             int compare = compareToNormal(i,testResult[i],testNormalLow[i],testNormalHigh[i]);
530
531
532             if(compare == COMPARE_TEST_LOW)
533             {
534                 haveAbnormalLowTests = true;
535
536                 abnormalLowTestExplanation.
537                 addMessage(ABNORMAL_LOW_TEST_EXPLANATION[i]);
538             }
539             else if(compare == COMPARE_TEST_HIGH)
540             {
541                 haveAbnormalHighTests = true;
542
543                 abnormalHighTestExplanation.
544                 addMessage(ABNORMAL_HIGH_TEST_EXPLANATION[i]);
545             }
546
547
548             if(haveAbnormalHighTests || haveAbnormalLowTests)
549                 haveAbnormalTests = true;
550
551             DialogTable dt = new DialogTable();
552             if(isVerbose())
553             {
554                 Phrase[] header = new Phrase[TEST_OUT_MAX];
555                 header[TEST_OUT_NAME] = new Phrase(headerNames[TEST_FIELD_NAME]);
556                 header[TEST_OUT_RESULT] = new Phrase(headerNames[TEST_FIELD_RESULT]);
557                 header[TEST_OUT_NORMAL_LOW] = new Phrase(headerNames[TEST_FIELD_NORMAL_LOW]);

```

```

558 header[TEST_OUT_NORMAL_HIGH] = new Phrase(headerNames[TEST_FIELD_NORMAL_HIGH]);
559 header[TEST_OUT_YOUR_RESULT] = new Phrase(TEST_COLUMN_YOUR_RESULT);
560 header[TEST_OUT_UNITS] = new Phrase(headerNames[TEST_FIELD_UNITS]);
561 header[TEST_OUT_REFERENCE] = new Phrase(TEST_COLUMN_REFERENCE);
562
563 dt.addHeader(header);
564 for(int i=0;i<testRows.length;i++)
565 {
566     Phrase[] row = new Phrase[TEST_OUT_MAX+1];
567
568     row[TEST_OUT_NAME] = new Phrase(testRows[i][TEST_FIELD_NAME]);
569
570
571     row[TEST_OUT_RESULT] = new Phrase(testResult[i]);
572
573     row[TEST_OUT_NORMAL_LOW] = new Phrase(testNormalLow[i]);
574
575     row[TEST_OUT_NORMAL_HIGH] = new Phrase(testNormalHigh[i]);
576
577     boolean isNormal = isTestNormal(i,testResult[i],testNormalLow[i],testNormalHigh[i]);
578
579     row[TEST_OUT_YOUR_RESULT] = new Phrase(isNormal);
580
581     row[TEST_OUT_UNITS] = new Phrase(testRows[i][TEST_FIELD_UNITS]);
582     Phrase ref = choseReferencePhrase(i,testResult[i],testNormalLow[i],testNormalHigh[i]);
583
584     row[TEST_OUT_REFERENCE] = ref;
585
586     dt.addRow(row);
587 }
588 getAcknowledgement(dt);

```

```

589     }
590     else
591     {
592         if(haveAbnormalTests)
593             getAcknowledgement(new Phrase("All your tests were normal with the exception of the following
594             tests."));
595         else
596             getAcknowledgement(new Phrase("All your tests were normal. There were no abnormal results."));
597
598         if(haveAbnormalTests)
599         {
600             // terse mode - no headers
601             for(int i=0;i<testRows.length;i++)
602             {
603                 boolean isNormal = isTestNormal(i,testResult[i],testNormalLow[i],testNormalHigh[i]);
604
605                 if(! isNormal)
606                 {
607                     Phrase[] row = new Phrase[TEST_TERSE_OUT_MAX+1];
608
609                     row[TEST_OUT_NAME] = new Phrase(testRows[i][TEST_FIELD_NAME]);
610
611                     row[TEST_OUT_RESULT] = new Phrase(testResult[i]);
612
613                     row[TEST_OUT_UNITS] = new Phrase(testRows[i][TEST_FIELD_UNITS]);
614
615                     dt.addRow(row);
616                 }
617             }
618         }
619     }

```

```

620         getAcknowledgement(dt);
621     }
622     getAcknowledgement(new Phrase(TEST_EXPLANATION));
623     if(haveAbnormalLowTests)
624         getAcknowledgement(abnormalLowTestExplanation);
625     if(haveAbnormalHighTests)
626         getAcknowledgement(abnormalHighTestExplanation);
627 }
628
629
630 private Phrase choseReferencePhrase(int testIndex, double testResult, double normalLow, double normalHigh)
631 {
632     String ref = "";
633
634     int compare = compareToNormal(testIndex, testResult, normalLow, normalHigh);
635
636     if(compare == COMPARE_TEST_NORMAL)
637         ref =
638         testRows[testIndex][TEST_FIELD_REFERENCE_NORMAL];
639     else if(compare == COMPARE_TEST_LOW)
640         ref = testRows[testIndex][TEST_FIELD_REFERENCE_LOW];
641     else if(compare == COMPARE_TEST_HIGH)
642         ref = testRows[testIndex][TEST_FIELD_REFERENCE_HIGH];
643     else if(compare == COMPARE_TEST_INDETERMINATE)
644         ref = testRows[testIndex][TEST_FIELD_REFERENCE_INDETERMINATE];
645
646     return new Phrase(ref);
647 }
648
649 private boolean isTestNormal(int testIndex, double testResult, double normalLow, double normalHigh)
650 {

```



```

651 int compare = compareToNormal(testIndex,testResult,normalLow,normalHigh);
652 if(compare == COMPARE_TEST_NORMAL)
653     return true;
654 return false;
655 }
656
657
658 private int compareToNormal(int testIndex,double testResult, double testNormalLow, double testNormalHigh)
659 {
660     int compare = 0;
661     if(testIndex == TEST_INDEX_ALKALINE_PHOSPHATASE)
662     {
663         compare = compareForAlkalinePhosphatase(age,sex,testResult);
664     }
665     else
666     {
667         if(testResult < testNormalLow)
668             compare = COMPARE_TEST_LOW;
669         if(testResult > testNormalHigh)
670             compare = COMPARE_TEST_HIGH;
671     }
672     //
673     return compare;
674 }
675
676
677 private static int REFTABLE_COL_AGE_LOW      = 0;
678 private static int REFTABLE_COL_AGE_HIGH     = 1;
679 private static int REFTABLE_COL_VALUE_LOW    = 2;
680 private static int REFTABLE_COL_VALUE_HIGH   = 3;
681

```

```

682 private int[][] alkalinePhosphataseRefValByAgeMale =
683 {
684     // low age, high age, lowNormal, highNormal
685     { 1, 9, 145, 420 },
686     { 10, 11, 130, 560 },
687     { 12, 13, 200, 495 },
688     { 14, 15, 130, 525 },
689     { 16, 19, 65, 260 }
690 };
691
692 private int[][] alkalinePhosphataseRefValByAgeFemale =
693 {
694     // low age, high age, lowNormal, highNormal
695     { 1, 9, 145, 420 },
696     { 10, 11, 130, 560 },
697     { 12, 13, 105, 420 },
698     { 14, 15, 70, 230 },
699     { 16, 19, 50, 130 }
700 };
701
702 // return 0 if normal, +1 for high, -1 for low and -2 for indeterminate
703 private int compareForAlkalinePhosphatase(int age,String sex,double testValue)
704 {
705     int[][] refTable = null;
706     if(sex.equalsIgnoreCase("male"))
707     {
708         refTable = alkalinePhosphataseRefValByAgeMale;
709     }
710     else
711     {
712         refTable = alkalinePhosphataseRefValByAgeFemale;

```

```

713     }
714
715     for(int i=0;i<refTable.length;i++)
716     {
717         if(age >= refTable[i][REFTABLE_COL_AGE_LOW] && age <=
718             refTable[i][REFTABLE_COL_AGE_HIGH])
719         {
720             if(testValue <
721                 refTable[i][REFTABLE_COL_VALUE_LOW])
722                 return COMPARE_TEST_LOW;
723             if(testValue >
724                 refTable[i][REFTABLE_COL_VALUE_HIGH])
725                 return COMPARE_TEST_HIGH;
726             return COMPARE_TEST_NORMAL; // must be normal
727         }
728     }
729
730     return COMPARE_TEST_INDETERMINATE; // not in the age range - indeterminate
731 }
732
733

```

2025040100

EXHIBIT 3

EXHIBIT 3

PSEUDO CODE LISTING OF TEST RESULT DELIVERY FOR STREPOCOCCUS TEST

```

1 package com.solveitech.st_notify;
2
3 import java.util.*;
4
5 import com.solveitech.st_lang.*;
6
7 public class StrepTest_Dialog extends DialogBase
8 {
9
10     private final static String ATTR_DIALOG_YOUR_STREP_RESULTS = "dialog.strep_results";
11     private final static String ATTR_DIALOG_STREP_REFERENCE = "dialog.strep_reference";
12     private final static String TEST_RESULT_MESSAGE_INTRO = "Your recent throat culture for Streptococcus was";
13
14     private static final String ABNORMAL_TEST_EXPLANATION =
15         "This germ can cause sore throat and fever and enlargement of neck glands. Even if you do not have these symptoms
16         you should receive an antibiotic to eradicate the germ so you do not have other complications.";
17
18     private static final String NORMAL_TEST_EXPLANATION =
19         "This test showed no evidence of Streptococcus. Other germs besides Strep can cause sore throat, fever and signs of
20         illness."
21
22     private static final String INDETERMINATE_TEST_EXPLANATION =
23         "The test for Strep could not be processed because of laboratory technical difficulties or may take somewhat longer to
24         process. If we are able to process it in spite of these difficulties, we will forward you the result in 24 hours.";
25
26     private boolean testResultPositive = false;

```

```

27 private String testResultReference = "";
28
29 //=====
30 public StrepTest_Dialog()
31 {
32
33     super();
34 }
35 final protected void init()
36     throws Exception
37 {
38     testResultPositive =
39     getAttrBoolean(ATTR_DIALOG_YOUR_STREP_RESULTS);
40     testResultReference = getAttrString(ATTR_DIALOG_STREP_REFERENCE);
41 }
42
43 final protected void register()
44 {
45
46     // registration is a way of telling scheduler what attributes are
47     // needed for a message.
48
49     registerAttribute(ATTR_DIALOG_YOUR_STREP_RESULTS);
50     registerAttribute(ATTR_DIALOG_STREP_REFERENCE);
51
52     // register the prompts that we are using. So that
53     // they can be verified before the conversation begins.
54
55
56     // register the roles that we require
57     registerRole(ROLE_PATIENT);

```

```

58     registerRole(ROLE_DOCTOR);
59 }
60
61 final protected void term()
62     throws Exception
63 {
64 }
65
66 final protected void execute()
67 {
68     // login has already been done.
69
70     Phrase testResultPhrase = new
71     Phrase(TEST_RESULT_MESSAGE_INTRO);
72
73     if(testResultPositive)
74         testResultPhrase.addMessage("POSITIVE");
75
76         getAcknowledgement(new Phrase(ABNORMAL_TEST_EXPLANATION);
77
78     if(testResultPositive == "NEGATIVE")
79         testResultPhrase.addMessage("NEGATIVE");
80         getAcknowledgement(new Phrase(NORMAL_TEST_EXPLANATION);
81
82     if(testResultPositive == "INDETERMINATE")
83         testResultPhrase.addMessage("INDETERMINATE");
84         getAcknowledgement(new Phrase(INDETERMINATE_TEST_EXPLANATION);
85
86         if(!getAcknowledgement(testResultPhrase))
87         {
88             setStatus(DIALOG_STATUS_FAIL);

```

```

89     return;
90 }
91
92 // for verbose mode, give the reference to Streptococcus
93 if(isVerbose())
94     getAcknowledgement(new Phrase(testResultReference));
95 }
96 }
97

```